

IN THE SPECIFICATION:

Page 39, second full paragraph, kindly replace with the following replacement paragraph:

ai The first strand of cDNA was then poly-G tailed using terminal transferase (Pharmacia, Freiburg) according to standard protocol. The tailed cDNA was PCR-amplified using a sense primer containing a poly-C stretch, based on the anchor primer sequence published by Gilliland, L. K. et al., (Tissue Antigens 47, 1-20, 1996) and designated 5'-AncTail (CGTCGATGAGCTCTAGAATTCCCCCCCCCCCCC[D]C, SEQ ID NO:7). This anchor primer was combined with an antisense primer, specific for the nucleotide sequence encoding the C-terminus of the lambda light chain constant region (CL2) or that of the IgG1-CH1 heavy chain domain (CGd1), respectively.

IN THE SPECIFICATION (cont.):

Page 40, Table 3, kindly replace with the following replacement Table 3:

Table 3: List of primers

5'-VH primer set:

VH1,3,5,7: AGGTGCAGCTGCTCGAGTCTGG (SEQ ID NO:8)
VH2: CAG(AG)TCACCTTGCTCGAGTCTGG (SEQ ID NO:9)
VH4: CAGGTGCAGCTGCTCGAGTCGGG (SEQ ID NO:10)
VH4B: CAGGTGCAGCTACTCGAGTGGGG (SEQ ID NO:11)
VH6: CAGGTACAGCTGCTCGAGTCAGG (SEQ ID NO:12)

3'-VH primer:

CGd1 GCATGTACTAGTTTTGTCACAAGATTG (SEQ ID NO:13)

5' VL primer set:

VL1: AATTTTGAGCTCACTCAGCCCCAC (SEQ ID NO:14)
VL2: TCTGCCGAGCTCCAGCCTGCCTCCGTG (SEQ ID NO:15)
VL3: TCTGTGGAGCTCCAGCCGCCCTCAGTG (SEQ ID NO:16)
VL4: TCTGAAGAGCTCCAGGACCCTGTTGTGTCTGTG (SEQ ID NO:17)
VL5: CAGTCTGAGCTCACGCAGCCGCCC (SEQ ID NO:18)
VL6: CAGACTGAGCTCACTCAGGAGCCC (SEQ ID NO:19)
VL7: CAGGTTGAGCTCACTCAACCGCCC (SEQ ID NO:20)
VL8: CAGGCTGAGCTCACTCAGCCGTCTTCC (SEQ IDNO:21)

3' VL primer:

CL2: CGCCGTCTAGAATTATGAACATTCTGTAGG (SEQ ID NO:22)